

SEQUENCE LISTING

<110> University of Rochester
Zauderer, Maurice
Ernest S. Smith

<120> In Vitro Methods Of Producing And Selecting
Immunoglobulin Molecules In Eukaryotic Cells

<130> 1821.0070004

<150> 60/271,424

<151> 2001-02-27

<150> 60/262,067

<151> 2001-01-18

<150> 60/298,087

<151> 2001-06-15

<150> 60/249,268

<151> 2000-11-17

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<170> PatentIn version 3.1

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Publ. No. 95423660

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<400> 96

aaagcggccg ccccgggatg ttacgtcc

28

Sequence

<210> 97
<211> 29
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<220>

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<400> 97
aaagggcccg gcgcgcctca ttgtttgcc

29

<210> 98
<211> 37
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<220>

<223> primer

<400> 98
aaaggatcca taatgaattc agtgactgta tcacacg

37

<210> 99
<211> 34
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<213> Artificial Sequence

<220>

<223> primer

<400> 99
cttgcgccg cttataaat aaaccottga gccc

34

<210> 100
<211> 34
<212> DNA
<213> Artificial Sequence

<220>

<223> primer

<400> 100

attgagctct taatactttt gtcgggtaac agag

34

<210> 101

<211> 29

<212> DNA

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<220>

<223> primer

<400> 101

ttactcgaga gtgtcgcaat ttggatttt

29

<210> 102

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 102

aaagaattcc tttattgtca tcggccaaa

29

<210> 103

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> primer

<400> 103

aatctgcagt cattgtttgc ctccctgctg

30

<210> 104

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 104
aaagaattca taatgaattc agtgactgta tcacacg

37

<210> 105

<211> 32

<212> DNA

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<220>

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<400> 105
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32

<210> 106

<211> 27

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<213> Artificial Sequence

<220>

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<400> 106
aataagcttt actccagata atatgga

27

<210> 107

<211> 23

<212> DNA

<213> Artificial Sequence

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<220>

<223> primer

<400> 107

aatctgcagc ccagttccat ttt

23

<210> 108

<211> 23

<212> DNA

<213> Artificial Sequence

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<223> primer

<400> 108

aatggatcct catccagcgg cta

23

<210> 109

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 109

aatgagctct agtacctaca acccgaa

27

<210> 110

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<223> primer

<400> 110

aaagtcgacg gccaaaaatt gaaatttt

28

Patent 2010/000000

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<211> 25

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<223> primer

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aatggatcct cattgtttgc ctccc

25

<210> 112

<211> 51

<212> DNA

<213> Artificial Sequence

<220>

<223> cassette converting Plasmid p7.5/tk3 to p7.5/tk3.1

<400> 112

gcggccgccc atggatagcg tgcacttgac tcgagaagct tagtagtcga c

51

<210> 113

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> region substituted to convert plasmid p7.5/tk3.1 to p7.5/tk3.2

<400> 113

ctcgagaagc ttagtagtcg ac

22

<210> 114

<211> 78

<212> DNA

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16

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60

81

<213> Artificial Sequence

42

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1 5 10

1

5

10

<210> 121

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 121

attagatcc ggtcaccgtc tcctcagcc

29

<210> 122

<211> 34

<212> DNA

<213> Artificial Sequence

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<223> primer

<400> 122

attagtcgac tcatttaccc ggagacaggg agag

34

<210> 123

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 123

aatatggta ccgtctcctc agcc

24

<210> 124

<211> 36

<212> DNA

<213> Artificial Sequence

<223> primer

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 $\langle 222 \rangle \quad (2) \dots (3)$

<223> May be any nucleotide

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<222> (5) . . (6)

<223> May be any nucleotide

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<222> (8) . . (9)

<223> May be any nucleotide

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 $\langle 222 \rangle \quad (11) \dots (12)$

<223> May be any nucleotide

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<223> May be any nucleotide

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$$\langle 222 \rangle \quad (17) \dots (18)$$

<223> May be any nucleotide

<400> 124

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36

<210> 125

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<220>

<221> misc_feature

<222> (1)..(2)

<223> May be any Nucleotide

<220>

<221> misc_feature

<222> (4)..(5)

<223> May be any Nucleotide

<220>

<221> misc_feature

<222> (7)..(8)

<223> May be any Nucleotide

<220>

<221> misc_feature

<222> (10)..(11)

<223> May be any Nucleotide

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<220>

<221> misc_feature

<222> (13)..(14)

<223> May be any Nucleotide

<220>

<221> misc_feature

<222> (16)..(17)

<223> May be any Nucleotide

<400> 125

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36

<210> 126

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 126

aatatgtcga ctcatttacc cgg

23

<210> 127

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 127

acacggtcac cgtctcctca gggagtgc

28

<210> 128

125-128

<213> Artificial Sequence

<223> primer

31

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30

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33

<213> Artificial Sequence

21

31

30

30

<210> 135

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 135

ttgttcaagg tgaaagtgaa gagaaaggaa

30

<210> 136

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 136

attagaattc atgcttgggg gtccagga

28

<210> 137

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 137

attagatcc tcacggcttc tccagctg

28

<210> 138

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 138

attaggatcc atggccaggc tggcggtg

28

<210> 139

<211> 34

<212> DNA

<213> Artificial Sequence

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<223> primer

<400> 139

attaccagca cactgggtcac tcctggcctg ggtg

34

<210> 140

<211> 69

<212> DNA

<213> Artificial Sequence

<220>

<223> p7.5/tk promoter

<220>

<221> CDS

<222> (46)..(69)

<223>

<400> 140

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Met Gly Pro Ala
1

57

gcc aac ggc gga
Ala Asn Gly Gly
5

69

<210> 141

<213> Artificial Sequence

<400> 141

<211> 75

<213> Artificial Sequence

<223> pE/Ltk promoter

<221> CDS

<222> (52) .. (75)

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                                     Met Gly  
                                     1
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ccg gcc gcc aac ggc gga 75
Pro Ala Ala Asn Gly Gly
5

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> tk sequence of pE/Ltk

<400> 143

Met Gly Pro Ala Ala Asn Gly Gly
1 5

<210> 144

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 144
aatatgcgcg cactcccagg tcaccttgaa ggagtctgg

39

<210> 145

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 145
aatatgcgcg cactccgagg tgcagctggt gcagtctg

38

<210> 146

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Signal Sequence

<400> 146
Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly

Ala His Ser

<211> 26

<212> PRT

<213> Artificial Sequence

 $\langle 220 \rangle$

<223> Signal Sequence

<400> 147

Asn Leu Trp Thr Thr Ala Ser Thr Phe Ile Val Leu Phe Leu Leu Ser
1 5 10 15

Leu Phe Tyr Ser Thr Thr Val Thr Leu Phe
20 25